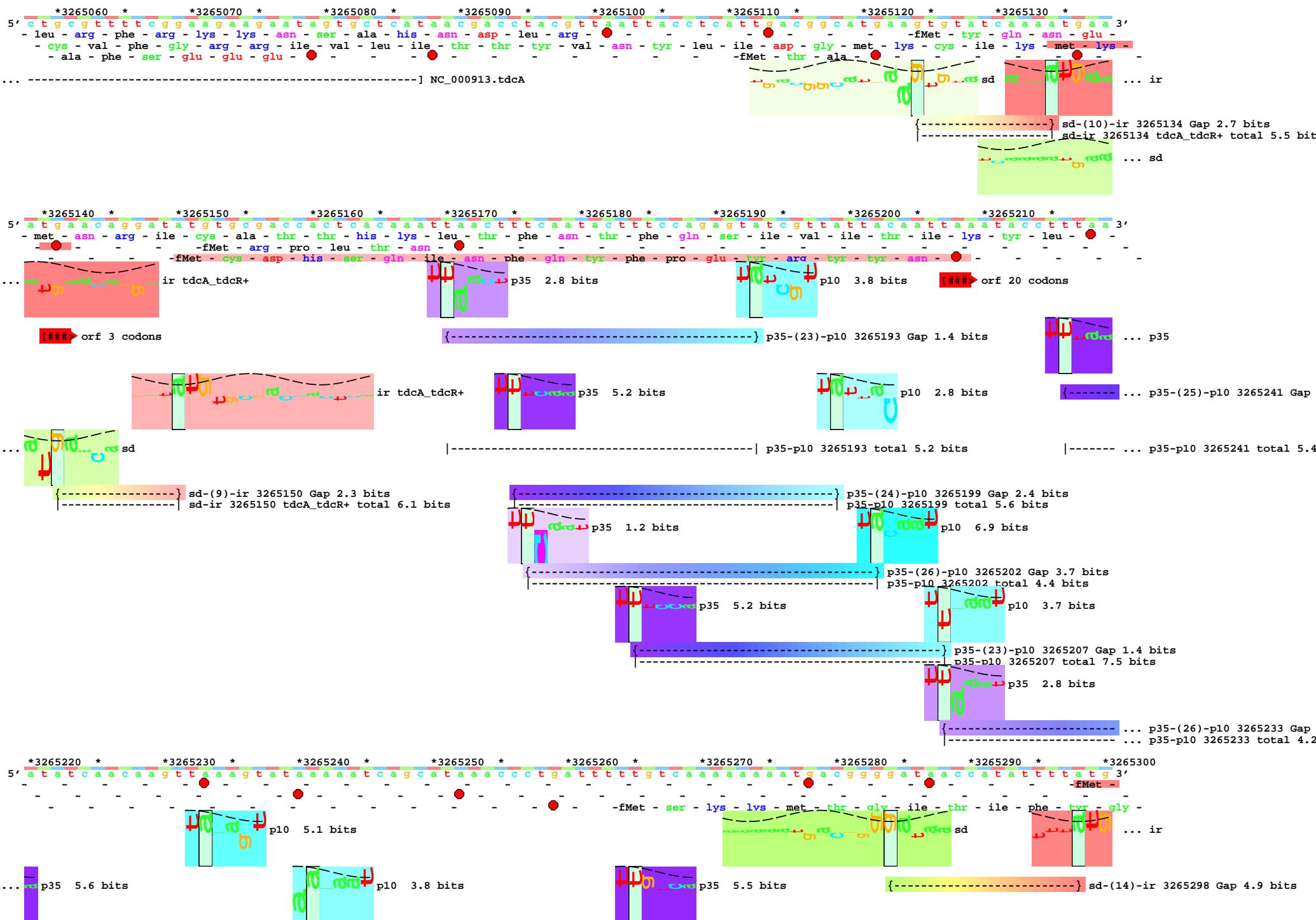


- 1 -

piece 1, NC_000913, tdcA_tdcR+, config: linear, direction: +, begin: 3265058, end: 3265421



```
... -----} p35-(25)-p10 3265241 Gap 4.0 bits  
... -----| p35-p10 3265241 total 5.4 bits
```

|-----| sd-ir 3265298 tdcA_tdcR+ total 8.6
 p10 1.1 bits

```
{ } p35-(23)-p10 3265288 Gap 1.4 bits  
|-----| p35-p10 3265288 total 5.2 bits
```

{ p35-(26)-p10 3265291 Gap 3.7 bits
| p35-p10 3265291 total 5.4 bits

```
... [REDACTED] p35-(26)-p10 3265233 Gap 3.7 bits  
... [REDACTED] p35-p10 3265233 total 4.2 bits
```

The diagram illustrates the tdcA gene structure. The top part shows the DNA sequence with transcription start sites indicated by asterisks (*). The bottom part shows the corresponding amino acid sequence with codons. A red box highlights the first 16 codons (ala-ile-thr-ser-phe-val-met-trp-leu-ile-gln-lys-arg-gly) which are part of the tdcA gene. A green box highlights the next 16 codons (asp-asn-ile-ile-arg-tyr-val-val-asn-thr-lys-lys-gly-leu-arg-pro-tyr-phe-lys-gln-leu-pro) which are part of the tdcR+ gene. A blue box highlights the final 16 codons (asp-asn-tyr-cln-alanine) which are part of the downstream gene. Red dots indicate transcription start sites for tdcA, and a red arrow indicates the direction of transcription for tdcR+. A green arrow indicates the direction of transcription for the downstream gene.

[###> orf 16 codons

```

{- ... sd-(7)-ir 3265388 Gap
|- ... sd-ir 3265388 tdcA_tdcR+
p35 2.3 bits

{----- ... p35-(24)-p10 3265399 Gap
----- ... p35-p10 3265399 total 4.3

```

The diagram illustrates a cDNA sequence with five start sites indicated by asterisks (*). The sequence is as follows:

```

5' - c a a a t t g a g t t a a t t t a a t g a g c a a g t t c t c a a a t t t 3'
      *   *3265390   *   *3265400   *   *3265410   *   *3265420
      -fMet- ser -    -    -    -    -    -    -    -    -    -    -    -    -
      lys - phe - glu - leu - asn - leu - met - ser - lys - phe - ser - asn -
  
```

Start sites are marked with red dots above the sequence. The first start site is at position 1, followed by positions 3265390, 3265400, 3265410, and 3265420. The amino acid sequence is shown below the sequence, starting with fMet at position 1.

[-----] ... NC 000913.tdcR

sd-(7)-ir 3265388 Gap 3.7 bits
sd-ir 3265388.tdcA tdcR+ total 5.6 bits

... -----} p35-(24)-p10 3265399 Gap 2.4 bits
-----| p35-p10 3265399 total 4.3 bits

[NH₂] orf 3 codons p10 4.5 bits